```
SEQUENCE LISTING
<110> Summers, Anne O.
     Caquiat, Jonathan
<120> Metal Binding Proteins, Recombinant Host Cells and
     Methods
<130> 79-00
<140> unassigned
<141> 2001-10-12
<150> US 60/240,465
<151> 2000-10-12
<160> 18
<170> PatentIn Ver. 2.0
<210> 1
<211> 435
<212> DNA
<213> Shigella flexneri, Tn21 of Plasmid R100
<400> 1
atggaaaata atttggaaaa cctgaccatt ggcgtttttg ccaaggcggc cggggtcaac 60
```

gtggagacaa tccgcttcta tcagcgcaag ggcctgttgc gggaaccgga caagccttac 120 ggcagcatcc gccgctatgg ggaggcggac gtggttcggg tgaaattcgt gaaatcggca 180 cagcggctgg ggttcagtct ggacgagatt gccgagctgt tgcggctcga cgatggcacc 240 cactgcgagg aggccagcag cctggccgaa cacaagctca aggacgtgcg cgagaagatg 300 geogaettgg egegeatgga aaccgtgetg tetgaacteg tgtgegeetg ceatgeacga 360 aaggggaatg tttcctgccc gttgatcgcg tcactacagg gcgaagcagg cctggcaagg 420 tcagctatgc cttag

```
<210> 2
<211> 144
<212> PRT
<213> Shigella flexneri, Tn21 of Plasmid R100
```

Met Glu Asn Asn Leu Glu Asn Leu Thr Ile Gly Val Phe Ala Lys Ala

Ala Gly Val Asn Val Glu Thr Ile Arg Phe Tyr Gln Arg Lys Gly Leu

Leu Arg Glu Pro Asp Lys Pro Tyr Gly Ser Ile Arg Arg Tyr Gly Glu 35

Ala Asp Val Val Arg Val Lys Phe Val Lys Ser Ala Gln Arg Leu Gly 55

Phe Ser Leu Asp Glu Ile Ala Glu Leu Leu Arg Leu Asp Asp Gly Thr 75 65

His Cys Glu Glu Ala Ser Ser Leu Ala Glu His Lys Leu Lys Asp Val 85 90 95

Arg Glu Lys Met Ala Asp Leu Ala Arg Met Glu Thr Val Leu Ser Glu
100 105 110

Leu Val Cys Ala Cys His Ala Arg Lys Gly Asn Val Ser Cys Pro Leu 115 120 125

Ile Ala Ser Leu Gln Gly Glu Ala Gly Leu Ala Arg Ser Ala Met Pro 130 135 140

<210> 3

<211> 321

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: chelon

<400> 3

atgacacact gcgaggaggc cagcagcctg gccgaacaca agctcaagga cgtgcgcgag 60 aagatggccg acttggcgcg catggaaacc gtgctgtctg aactcgtgtg cgcctgccat 120 gcacgaaagg ggaatgtttc ctgcccgttg atcgcgtcac tacagggatc ctcaggcacc 180 gccgacttgg aggccatgga caccgtgctg tctgaactcg tgtgcgcctg ccatgcacga 300 aaggggaatg tttcctgcc g

<210> 4

<211> 117

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: chelon

<400> 4

Met Thr His Cys Glu Glu Ala Ser Ser Leu Ala Glu His Lys Leu Lys

1 5 10 15

Asp Val Arg Glu Lys Met Ala Asp Leu Ala Arg Met Glu Thr Val Leu 20 25 30

Ser Glu Leu Val Cys Ala Cys His Ala Arg Lys Gly Asn Val Ser Cys 35 40 45

Pro Leu Ile Ala Ser Leu Gln Gly Ser Ser Gly Thr His Cys Glu Glu
50 55 60

Ala Ser Ser Leu Ala Glu His Lys Leu Lys Asp Val Arg Glu Lys Met 65 70 75 80

Ala Asp Leu Ala Arg Met Glu Thr Val Leu Ser Glu Leu Val Cys Ala 85 90 95

Cys His Ala Arg Lys Gly Asn Val Ser Cys Pro Ser Ala Trp Ser His 100 105 110

Pro Gln Phe Glu Lys 115

<210> 5

<211> 117

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: chelon

<400> 5

Met Thr His Cys Glu Glu Val Ser Ser Leu Ala Glu His Lys Leu Lys

1 5 10 15

Asp Val Arg Glu Lys Met Ala Asp Leu Ala Arg Met Glu Thr Val Leu 20 25 30

Ser Glu Leu Val Cys Ala Cys His Ala Arg Lys Gly Asn Val Ser Cys
35 40 45

Pro Leu Ile Ala Ser Leu Gln Gly Ser Ser Gly Thr His Cys Glu Glu 50 55 60

Val Ser Ser Leu Ala Glu His Lys Leu Lys Asp Val Arg Glu Lys Met 65 70 75 80

Ala Asp Leu Ala Arg Met Glu Thr Val Leu Ser Glu Leu Val Cys Ala 85 90 95

Cys His Ala Arg Lys Gly Asn Val Ser Cys Pro Ser Ala Trp Ser His
100 105 110

Pro Gln Phe Glu Lys 115

<210> 6

<211> 118

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: chelon

<400> 6

Met Thr His Cys Glu Glu Ala Ser Ser Leu Val Glu His Lys Leu Lys

1 5 10 15